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SEQUENCE LISTING

<110> Autogen Research Pty Ltd

<120> Novel genes and their use in the modulation of obesity, diabetes and energy imbalance

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<151> 1999-06-29

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<170> PatentIn Ver. 2.0

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aga tgc acc gtt gta aca cca aaa gtc ttc cga gtc gga gaa tat gaa      198
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caa gtt aca ttt gaa gcc cac ggt cac act gac cca ttt gat gta acc      246
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Ser Val His Leu Ser Pro Glu Asn Lys Phe Lys Asn Ser Thr Ile Leu	
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Leu Glu Arg Gln Arg Cys Thr Val Val Thr Pro Lys Val Phe Arg Val

- 4 -

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Phe Asp Val Thr Ile Ser Ile Lys Ser Tyr Pro Asp Lys Asn Ala Asn		
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Tyr Ser Ser Ser Ser Val His Leu Ser Pro Glu Asn Lys Phe Lys Asn		
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Ser Thr Ile Leu Thr Ile Gln Pro Lys Gln Leu Ser Glu Gly Gln Asn		
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Ser Ser Ser His Val Tyr Leu Glu Val Val Ser Lys His Phe Ser Thr		
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Ser Lys Ile Met Ser Ile Val Tyr Asp Asn Gly Thr Leu Phe Ile Gln		
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Tyr Ser Leu Asp Glu Ala Leu Lys Pro Val Thr Arg Glu Thr Val Leu		
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Asn His Thr Gly Ile Thr Ser Phe Pro Asp Phe Arg Ile Pro Thr Asn		
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Pro Lys Pro Gly Arg Trp Met Ile Lys Ala Lys Tyr Arg Glu Asp Ala		
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Phe Lys Ile Ala Leu Val Pro Thr Ser Asp Leu Glu His Pro Met Glu		
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Glu Ala Arg Gly Leu Ser Leu Gln Pro Lys Lys Ser Leu Gln Glu Met		
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Ile His Glu Gln Ala Ser Lys Tyr Lys His Pro Val Leu Lys Lys Cys		
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Cys Tyr Asp Gly Ala Arg Tyr Asn His His Glu Thr Cys Glu Glu Arg		
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Val Ala Arg Val Lys Ile Gly Pro Asn Cys Val Arg Ala Phe Ser Glu		
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		320

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 Pro Ala Leu Glu Thr Glu Gly Leu Arg Phe Leu His Val Thr Val Gly
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tcc ctg ctg gcc agc tat ggc tgg tac gtc ctc ttc agc tgc atc ctt 148
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 Leu Tyr Ile Val Ile Gln Lys Leu Ser Val Arg Leu Arg Val Leu Arg
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cag agg cag ctg gac cag gct gac gct gtt ctg gaa cct gat gct gtt 244
 Gln Arg Gln Leu Asp Gln Ala Asp Ala Val Leu Glu Pro Asp Ala Val
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Asp Lys Lys Pro Leu Arg Gly Asn Gly Tyr Asn Pro Leu Thr Gly Glu
160 165 170

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Gln Lys Leu Ser Val Arg Leu Arg Val Leu Arg Gln Arg Gln Leu Asp
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Gln Ala Asp Ala Val Leu Glu Pro Asp Ala Val Val Lys Arg Gln Glu
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Ala Leu Ala Ala Ala Arg Leu Arg Met Gln Glu Asp Leu Asn Ala Gln
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Val Glu Lys His Lys Glu Lys Leu Arg Gln Leu Glu Glu Glu Lys Arg
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Arg Gln Lys Ile Glu Met Trp Asp Ser Met Gln Glu Gly Arg Ser Tyr
      115          120          125

Arg Arg Asn Pro Gly Arg Pro Gln Glu Glu Asp Gly Pro Gly Pro Ser
      130          135          140

Thr Ser Ser Ser Val Thr Arg Lys Gly Lys Ser Asp Lys Lys Pro Leu
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Ala Trp Arg Pro Gly Arg Arg Gly Pro Ser Ser Gly Gly
      180          185
  
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 Ser Gly Gly
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Gln Lys Leu Ser Ala Arg Leu Arg Ala Leu Arg Gln Arg Gln Leu Asp
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Arg Ala Ala Ala Ala Val Glu Pro Asp Val Val Val Lys Arg Gln Glu
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Ala Leu Ala Ala Ala Arg Leu Lys Met Gln Glu Glu Leu Asn Ala Gln
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Val Glu Lys His Lys Glu Lys Leu Lys Gln Leu Glu Glu Glu Lys Arg

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100              105              110

Arg Gln Lys Ile Glu Met Trp Asp Ser Met Gln Glu Gly Lys Ser Tyr
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Lys Gly Asn Ala Lys Lys Pro Gln Glu Glu Asp Ser Pro Gly Pro Ser
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Thr Ser Ser Val Leu Lys Arg Lys Ser Asp Arg Lys Pro Leu Arg Gly
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Thr Val Phe Leu Leu Val Thr Ser Leu Pro His
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